#2

PAGE: 1

RAW SEQUENCE LISTING PATENT APPLICATION US/08/086,427

DATE: 08/01/93 TIME: 12:20:01

1		SEQUENCE LISTING
2	(1) 0	anoual Information
3 4	(1) G	eneral Information: APPLICANT: Gospodarowicz, Denis
5 6 7	(i)	APPLICANT: Gospodarowicz, Denis Masiarz, Frank R.
8 9	(ii)	TITLE OF INVENTION: A Truncated Keratinocyte Growth Factor (KGF) Having Increased Biological Activity
10	(222)	WHATER OF GEOVERNOES 10
11 12	(111)	NUMBER OF SEQUENCES: 19
13	(iv)	CORRESPONDENCE ADDRESS:
14	(10)	(A) ADDRESSEE: Chiron Corporation
15		(B) STREET: 4560 Horton Street
16		(C) CITY: Emeryville
17		(D) STATE: CA
18.		(E) COUNTRY: USA
19		(F) ZIP: 94608
20		
21	(v)	COMPUTER READABLE FORM:
22		(A) MEDIUM TYPE: Floppy disk
23		(B) COMPUTER: IBM PC compatible
24		(C) OPERATING SYSTEM: PC-DOS/MS-DOS
25		(D) SOFTWARE: PatentIn Release #1.0, Version #1.25
26		
27	(vi)	CURRENT APPLICATION DATA:
28		(A) APPLICATION NUMBER:
29		(B) FILING DATE:
30		(C) CLASSIFICATION:
31 32	(3ri i i)	ATTORNEY/AGENT INFORMATION:
33	(\ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \	(A) NAME: Amy L. Collins, Esq.
34		(B) REGISTRATION NUMBER: 33,370
35		(C) REFERENCE/DOCKET NUMBER: 0953.001
36		(c) Reference, booker normality opposite
37	(ix)	TELECOMMUNICATION INFORMATION:
38	(===,	(A) TELEPHONE: (510) 601-2768
39		(B) TELEFAX: (510) 655-3542
40		
41		
42	(2) INFO	RMATION FOR SEQ ID NO:1:
43.		•
44	(i)	SEQUENCE CHARACTERISTICS:
45		(A) LENGTH: 194 amino acids
46		(B) TYPE: amino acid
47		(C) STRANDEDNESS: single
48		(D) TOPOLOGY: linear
49	1225	NOT ECT E TWO TO THE TOTAL TO
50	(11)	MOLECULE TYPE: protein

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52																	
53																	
54		(xi)	SEQ	JENC	E DE	SCRI	PTIO	N: SI	EQ II	ON C	:1:						
55																	
56		Met	His	Lys	Trp	Ile	Leu	Thr	Trp	Ile	Leu	Pro	Thr	Leu	Leu	Tyr	Arg
57		1		_	_	5			_		10					15	_
58																	
59		Ser	Cys	Phe	His	Ile	Ile	Cys	Leu	Val	Gly	Thr	Ile	Ser	Leu	Ala	Cys
60 ·			-		20			-		25	•				30		-
61																	
62		Asn	Asp	Met	Thr	Pro	Glu	Gln	Met	Ala	Thr	Asn	Val	Asn	Cvs	Ser	Ser
63			-	35					40					45	-		
64																	
65		Pro	Glu	Ara	His	Thr	Ara	Ser	Tvr	Asp	Tvr	Met	Glu	Glv	Glv	Asp	Ile
66			50					55			-4		60	- 1			
67																	
68		Ara	Val	Ara	Ara	Leu	Phe	Cvs	Ara	Thr	Gln	Trp	Tvr	Leu	Ara	Ile	Asp
69		65		5	5		70	-1-	5			75	- 2		3		80
70												. –					
71		Lvs	Ara	Glv	Lvs	Val	Lvs	Glv	Thr	Gln	Glu	Met	Lvs	Asn	Asn	Tyr	Asn
72			3	2		85		2			90		- 4			95	
73																	
74		Ile	Met	Glu	Ile	Arq	Thr	Val	Ala	۷al	Gly	Ile	Val	Ala	Ile	Lys	Gly
75					100	•				105	•				110	-	-
76																	
77		Val	Glu	Ser	Glu	Phe	Tyr	Leu	Ala	Met	Asn	Lys	Glu	Gly	Lys	Leu	Tyr
78				115			-		120			•		125	-		-
79																	
80		Ala	Lys	Lys	Glu	Cys	Asn	Glu	Asp	Cys	Asn	Phe	Lys	Glu	Leu	Ile	Leu
81			130	•		•		135	•	•			140				
82																	
83		Glu	Asn	His	Tyr	Asn	Thr	Tyr	Ala	Ser	Ala	Lys	Trp	Thr	His	Asn	Gly
84		145			-		150	•				155	-				160
85																	
86		Gly	Glu	Met	Phe	Val	Ala	Leu	Asn	Gln	Lys	Gly	Ile	Pro	Val	Arg	Gly
87		-				165					170	-				175	_
88																	
89		Lys	Lys	Thr	Lys	Lys	Glu	Gln	Lys	Thr	Ala	His	Phe	Leu	Pro	Met	Ala
90		_	-		180	-			_	185					190		
91																	
92		Ile	Thr														
93																	
94																	
95	(2)	INFO	RMAT:	ION I	FOR S	SEQ :	ID NO	0:2:									
96																	
97		(i)	SEQ	JENCI	E CHA	ARAC'	rer is	STICS	3:								
98			(A)	LEI	NGTH	: 23	amir	no a	cids								
99						amino											
100			(C)	ST	RANDI	EDNE	SS: 8	sing.	le								
101			(D)	TO	POLO	GY: 3	linea	ar									
102																	

RAW SEQUENCE LISTING PATENT APPLICATION US/08/086,427

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100	(ii) MOLEGUE WARE washed
103 104	(ii) MOLECULE TYPE: protein
105	
106	
107	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
108	
109	Cys Asn Asp Met Thr Pro Glu Gln Met Ala Thr Asn Val Asn Cys Ser
110 111	1 5 10 15
112	Ser Pro Glu Arg His Thr Arg
113	20
114	
115	(2) INFORMATION FOR SEQ ID NO:3:
116	
117	(i) SEQUENCE CHARACTERISTICS:
118	(A) LENGTH: 6 amino acids
119	(B) TYPE: amino acid
120 121	(C) STRANDEDNESS: single (D) TOPOLOGY: linear
122	(b) Toronogi. Timear
123	(ii) MOLECULE TYPE: peptide
124	
125	
126	
127	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:
128	Mah Win Toon Man Tile Too
129 130	Met His Lys Trp Ile Leu 1 5
131	1 3
132	(2) INFORMATION FOR SEQ ID NO:4:
133	•
134	(i) SEQUENCE CHARACTERISTICS:
135	(A) LENGTH: 35 base pairs
136	(B) TYPE: nucleic acid
137 138	(C) STRANDEDNESS: single (D) TOPOLOGY: linear
139	(b) Topologi: Timear
140	(ii) MOLECULE TYPE: DNA (genomic)
141	(, (.
142	
143	
144	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:
145 146	AGATCTCTGC AGCTATAATG CACAAATGGA TACTG 35
146	MUNICICIAC MUCINIANIA CACAMAIAGA IACIG
148	(2) INFORMATION FOR SEQ ID NO:5:
149	(-, -::-:
150	(i) SEQUENCE CHARACTERISTICS:
151	(A) LENGTH: 7 amino acids
152	(B) TYPE: amino acid
153	(C) STRANDEDNESS: single

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154 155			(D) TOPOLOGY: linear
156		(ii)	MOLECULE TYPE: peptide
157		(,	
158			
159			
160		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:5:
161			
162		Thr	Ile Ala Met Pro Leu Phe
163		1	5
164			
165	(2)	INFO	RMATION FOR SEQ ID NO:6:
166			
167		(i)	SEQUENCE CHARACTERISTICS:
168			(A) LENGTH: 38 base pairs
169			(B) TYPE: nucleic acid
170			(C) STRANDEDNESS: single
171			(D) TOPOLOGY: linear
172			
173		(ii)	MOLECULE TYPE: DNA (genomic)
174			
175			
176			
177		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:6:
178			
179	AGA'.	rcrgc	GG CCGCTTAAGT TATTGCCATA GGAAGAAA 38
180	(0)	T1700	DVARTOV DOD GEO ID VO G
181	(2)	INFO	RMATION FOR SEQ ID NO:7:
182 183		(-)	CROUPLIOR CUARACTERICO.
184		(_ /	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 amino acids
185			(B) TYPE: amino acid
186			(C) STRANDEDNESS: single
187			(D) TOPOLOGY: linear
188			(b) TOFOLOGI: Tillear
189		(ii)	MOLECULE TYPE: peptide
190		(11)	Mondeodd III. peperae
191			
192			
193		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:7:
194		,,	
195		Ser	Tyr Asp Tyr Met Glu Gly Gly Asp Ile Arg Val Arg Arg Leu Phe
196		1	5 10 15
197			
198		Xaa	Arg Thr Gln
199			20
200			
201	(2)	INFO	RMATION FOR SEQ ID NO:8:
202			
203		(i)	SEQUENCE CHARACTERISTICS:
204			(A) LENGTH: 32 base pairs

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205	(B) TYPE: nucleic acid	
206	(C) STRANDEDNESS: single	
207	(D) TOPOLOGY: linear	
208		
209	(ii) MOLECULE TYPE: DNA (genomic)	
210.		
211		
212		
213	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:	
214		
215	GGTGGTGGAT CCCCAGCTTA GTTCATAGGT CC	32
216		
217	(2) INFORMATION FOR SEQ ID NO:9:	
218		
219	(i) SEQUENCE CHARACTERISTICS:	
220	(A) LENGTH: 12 amino acids	
221	(B) TYPE: amino acid	
222	(C) STRANDEDNESS: single	
223	(D) TOPOLOGY: linear	
224		
225	(ii) MOLECULE TYPE: peptide	
226	• •	
227		
228		
229	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:	
230	,	
	His Gln Asn Val Phe Arg Lvs Ala Pro Ile Gln Ala	
231	His Gln Asn Val Phe Arg Lys Ala Pro Ile Gln Ala	
231 232	His Gln Asn Val Phe Arg Lys Ala Pro Ile Gln Ala 1 5 10	
231 232 233	1 5 10	
231 232 233 234		
231 232 233 234 235	1 5 10 (2) INFORMATION FOR SEQ ID NO:10:	
231 232 233 234 235 236	1 5 10 (2) INFORMATION FOR SEQ ID NO:10: (i) SEQUENCE CHARACTERISTICS:	
231 232 233 234 235 236 237	1 5 10 (2) INFORMATION FOR SEQ ID NO:10: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs	
231 232 233 234 235 236 237 238	1 5 10 (2) INFORMATION FOR SEQ ID NO:10: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid	
231 232 233 234 235 236 237 238 239	1 5 10 (2) INFORMATION FOR SEQ ID NO:10: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single	
231 232 233 234 235 236 237 238 239 240	1 5 10 (2) INFORMATION FOR SEQ ID NO:10: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid	
231 232 233 234 235 236 237 238 239 240 241	1 5 10 (2) INFORMATION FOR SEQ ID NO:10: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
231 232 233 234 235 236 237 238 239 240 241 242	1 5 10 (2) INFORMATION FOR SEQ ID NO:10: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single	
231 232 233 234 235 236 237 238 239 240 241 242 243	1 5 10 (2) INFORMATION FOR SEQ ID NO:10: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
231 232 233 234 235 236 237 238 239 240 241 242 243 244	1 5 10 (2) INFORMATION FOR SEQ ID NO:10: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
231 232 233 234 235 236 237 238 239 240 241 242 243 244 245	1 5 10 (2) INFORMATION FOR SEQ ID NO:10: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic)	
231 232 233 234 235 236 237 238 239 240 241 242 243 244 245 246	1 5 10 (2) INFORMATION FOR SEQ ID NO:10: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
231 232 233 234 235 236 237 238 239 240 241 242 243 244 245 246 247	1 5 10 (2) INFORMATION FOR SEQ ID NO:10: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:	36
231 232 233 234 235 236 237 238 239 240 241 242 243 244 245 246 247 248	1 5 10 (2) INFORMATION FOR SEQ ID NO:10: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic)	36
231 232 233 234 235 236 237 238 239 240 241 242 243 244 245 246 247 248 249	1 5 10 (2) INFORMATION FOR SEQ ID NO:10: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: GTGTTGGTTA ACGAATCGCT TAGCCGGAAT TTGTGC	36
231 232 233 234 235 236 237 238 239 240 241 242 243 244 245 246 247 248 249 250	1 5 10 (2) INFORMATION FOR SEQ ID NO:10: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:	36
231 232 233 234 235 236 237 238 239 240 241 242 243 244 245 246 247 248 249 250 251	1 5 10 (2) INFORMATION FOR SEQ ID NO:10: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: GTGTTGGTTA ACGAATCGCT TAGCCGGAAT TTGTGC (2) INFORMATION FOR SEQ ID NO:11:	36
231 232 233 234 235 236 237 238 239 240 241 242 243 244 245 246 247 248 249 250 251 252	1 5 10 (2) INFORMATION FOR SEQ ID NO:10: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: GTGTTGGTTA ACGAATCGCT TAGCCGGAAT TTGTGC (2) INFORMATION FOR SEQ ID NO:11: (i) SEQUENCE CHARACTERISTICS:	36
231 232 233 234 235 236 237 238 239 240 241 242 243 244 245 246 247 248 249 250 251 252 253	1 5 10 (2) INFORMATION FOR SEQ ID NO:10: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: GTGTTGGTTA ACGAATCGCT TAGCCGGAAT TTGTGC (2) INFORMATION FOR SEQ ID NO:11: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 12 amino acids	36
231 232 233 234 235 236 237 238 239 240 241 242 243 244 245 246 247 248 249 250 251 252	1 5 10 (2) INFORMATION FOR SEQ ID NO:10: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: GTGTTGGTTA ACGAATCGCT TAGCCGGAAT TTGTGC (2) INFORMATION FOR SEQ ID NO:11: (i) SEQUENCE CHARACTERISTICS:	36

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256	(D) TOPOLOGY: linear	
257 258	(ii) MOLECULE TYPE: peptide	
258 259	(II) MODECODE TYPE: pepcide	
260		
261		
262	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:	
263		
264	Pro Ala Lys Arg Ser Tyr Asp Tyr Met Glu Gly Gly	
265	1 5 10	
266	·	
	(2) INFORMATION FOR SEQ ID NO:12:	
268		
269	(i) SEQUENCE CHARACTERISTICS:	
270	(A) LENGTH: 39 base pairs	
271	(B) TYPE: nucleic acid	
272	(C) STRANDEDNESS: single	
273 274	(D) TOPOLOGY: linear	
274	(ii) MOLECULE TYPE: DNA (genomic)	
276	(II) MODECODE TIFE. DNA (Genomic)	
277		
278		
279	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:	
280		
281 (CCGCCGGCTA AGCGAAGTTA TGATTACATG GAAGGAGGG	39
282		
	(2) INFORMATION FOR SEQ ID NO:13:	
284		
285	(i) SEQUENCE CHARACTERISTICS:	
286	(A) LENGTH: 8 amino acids	
287 288	(B) TYPE: amino acid (C) STRANDEDNESS: single	
289	(D) TOPOLOGY: linear	
290	(b) TOPOLOGI. Timeat	
291	(ii) MOLECULE TYPE: peptide	
292	(,	
293		
294		
295	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:	
296		
297	Thr Ile Ala Met Pro Leu Phe His	
298	1 5	
299	(a) TYPODYAMION FOR CEO ID NO 14	
	(2) INFORMATION FOR SEQ ID NO:14:	
301 302		
3 ∪⊿	(i) CROHENCE CUADACTEDICTICS.	
	(i) SEQUENCE CHARACTERISTICS:	
303	(A) LENGTH: 39 base pairs	

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307				
308		(ii)	MOLECULE TYPE: DNA (genomic)	
309				
310				
311				
312		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:14:	
313		(,		
314	сстс	стст	CG ACTTAAGTTA TTGCCATAGG AAGAAAGTG	39
315	0010	.0101	CO ACITAGITA TICCONTACO AMORAGIO	"
316	(2)	TNEO	RMATION FOR SEQ ID NO:15:	
317	(2)	INFO	RUMITON FOR BEQ ID NO.15.	
318		(i)	SEQUENCE CHARACTERISTICS:	
319		(1)	(A) LENGTH: 25 base pairs	
320			(B) TYPE: nucleic acid	
321			(C) STRANDEDNESS: single	
			(D) TOPOLOGY: linear	
322 323			(D) TOPOLOGY: Timear	
		1221	MOLEGIE EVER DIV (comonic)	
324		(11)	MOLECULE TYPE: DNA (genomic)	
325				
326				
327		1 .23	ABOTEMAE DEGARDERION AND TO NO 15	
328		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:15:	
329	~~		OT 122 TETTOGGG 02 TGG	٥.
330	GATO	CAGAT	CT AAATTTCCCG GATCC	25
331	(0)		NAME OF THE OFF THE TO THE	
332	(2)	INFO	RMATION FOR SEQ ID NO:16:	
333			A-A	
334		(1)	SEQUENCE CHARACTERISTICS:	
335			(A) LENGTH: 25 base pairs	
336			(B) TYPE: nucleic acid	
337			(C) STRANDEDNESS: single	
338			(D) TOPOLOGY: linear	
339				
340		(11)	MOLECULE TYPE: DNA (genomic)	
341				
342				
343				
344		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:16:	
345				
346	TCT	AGATT'	TA AAGGGCCTAG GAGCT	25
347				
348	(2)	INFO	RMATION FOR SEQ ID NO:17:	
349				
350		(i)	SEQUENCE CHARACTERISTICS:	
351			(A) LENGTH: 9 amino acids	
352			(B) TYPE: amino acid	
353			(C) STRANDEDNESS: single	
354			(D) TOPOLOGY: linear	
355				
356		(ii)	MOLECULE TYPE: DNA (genomic)	
357				

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358				
359				
360		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:17:	
361				
362			Ser Tyr Asp Tyr Met Glu Gly Gly	
363		1	5	
364	(0)	******	NATION TOD OTO TO WO 40	
365	(2)	INFO	RMATION FOR SEQ ID NO:18:	
366		(=)		
367		(1)	SEQUENCE CHARACTERISTICS:	
368			(A) LENGTH: 35 base pairs	
369			(B) TYPE: nucleic acid	
370			(C) STRANDEDNESS: single	
371			(D) TOPOLOGY: linear	
372			MOT TOTAL TOTAL PARTY ()	
373		(11)	MOLECULE TYPE: DNA (genomic)	
374				
375				
376 377		/ \	CROURNOE DECODIDATON. CEO ID NO.10.	
378		(XI)	SEQUENCE DESCRIPTION: SEQ ID NO:18:	
379	Cmm/	ammma:	AT GAGTTATGAT TACATGGAAG GAGGG	35
380	GII	3 111C	AI GAGIIAIGAI IACAIGGAAG GAGGG	33
381				
382	(2)	TMEO	PMARTON FOR CEO ID NO.10.	
383	(2)	INFO	RMATION FOR SEQ ID NO:19:	
384		(3)	SEQUENCE CHARACTERISTICS:	
385		(1)	(A) LENGTH: 99 base pairs	
386			(B) TYPE: nucleic acid	
387			(C) STRANDEDNESS: single	
388			(D) TOPOLOGY: linear	
389			(b) 10102001. 1211002	
390		(ii)	MOLECULE TYPE: DNA (genomic)	
391		(/	(30)	
392				
393		(ix)	FEATURE:	
394		((A) NAME/KEY: -	
395			(B) LOCATION: 1314	
396			(D) OTHER INFORMATION: /note= "The figure did not conta	in
397			the intervening polyhedrin sequences."	
398			3 L · 1	
399		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:19:	
400		,	-	
401	TAT	AAATA	TT CCGGGCGCGG ATCGGTACCA GATCTGCAGA ATTCTAGAGG	
402	ATC	CTGAT	CA 60	
403	_			
404	GCT	AGCAG	AG CTCGCGGCCG CCCGGGCCGT ACCGACTCT	99
405				

SEQUENCE VERIFICATION REPORT PATENT APPLICATION US/08/086,427

DATE: 08/01/93 TIME: 12:20:35

INPUT SET: S6238.raw

Line Error Original Text

29 399

Wrong Filing Date Entered (99) and Calc. Seq. Length (49) differ

(B) FILING DATE: (xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:

SEQUENCE MISSING ITEM REPORT PATENT APPLICATION US/08/086,427

DATE: 08/01/93 TIME: 12:20:35

INPUT SET: S6238.raw

APPLICATION NUMBER FILING DATE PRIOR APPLICATION DATA

SEQUENCE CORRECTION REPORT PATENT APPLICATION US/08/086,427

DATE: 08/01/93 TIME: 12:20:36

INPUT SET: S6238.raw

Line

Original Text

Corrected Text